



PCT

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/088,187A

DATE: 04/15/2003  
TIME: 13:52:40

#12

Input Set : A:\p02825.app  
Output Set: N:\CRF4\04152003\J088187A.raw

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3 41100 APPLICANT: Dean, Caroline
4      Levy, Yaron Y
5 41200 TITLE OF INVENTION: Methods and Means for Modification of Plant Flowering
6      Characteristics
7 41300 FILE REFERENCE: 0380-P02825US0
11 41400 CURRENT APPLICATION NUMBER: US 10/088,187A
12 41410 CURRENT FILING DATE: 2002-03-15
14 41500 PRIOR APPLICATION NUMBER: PCT/GB00/03525
15 41510 PRIOR FILING DATE: 2000-09-13
17 41500 PRIOR APPLICATION NUMBER: GB 9922071.7
18 41510 PRIOR FILING DATE: 1999-09-17
20 41600 NUMBER OF SEQ ID NOS: 48
21 41700 SOFTWARE: PatentIn Ver. 2.1
24 42100 SEQ ID NO: 1
25 42110 LENGTH: 5000
26 42120 TYPE: DNA
27 42130 ORGANISM: Arabidopsis thaliana
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104 acaactaaaa aactctcttc tgggtctgtt cattattgag tcagtgtctc gttttttctc 4500
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117 <213> ORGANISM: Arabidopsis thaliana
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177 <213> ORGANISM: Arabidopsis thaliana
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184 1 5
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186 <211> LENGTH: 4
187 <212> TYPE: PRT
188 <213> ORGANISM: Arabidopsis thaliana
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191 1 5
192 <210> SEQ ID NO: 8
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194 <212> TYPE: DNA
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197 <223> OTHER INFORMATION: Description of Artificial Sequence: vrnl-2
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200 <221> NAME/KEY: CDS
201 <222> LOCATION: (1)..(27)
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204 Lys Lys Met Leu Ile Leu Arg Lys
205 1 5
206 <210> SEQ ID NO: 9
207 <211> LENGTH: 9
208 <212> TYPE: PRT
209 <213> ORGANISM: Artificial Sequence
210 <220> FEATURE:
211 <223> OTHER INFORMATION: Description of Artificial Sequence: vrnl-2
212 mutation
213 <400> SEQUENCE: 9
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216 <210> SEQ ID NO: 10
217 <211> LENGTH: 1495
218 <212> TYPE: DNA
219 <213> ORGANISM: Arabidopsis thaliana

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Input Set : A:\p02825.app

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243 22211 NAME/KEY: CDS

244 22221 LOCATION: (769)..(1294)

246 24001 SEQUENCE: 10

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251 actctttcag atctgaaagg gttctttctt ctctttagtc tctttgtcct ttattttctc 180
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255 tcttacttgg tcaatgttgt gtctgaac atg cca cgc cct ttc ttc cat aag 300
256                                     Met Pro Arg Pro Phe Phe His Lys
257                                     1               5
258 ttg att ttc tca tcc act atc caa gaa aaa cgt ctg agg gtc cca gat 360
260 leu ile phe ser ser thr ile gln glu lys arg leu arg val pro asp
261 10               15               20
262 aag ttt gtg agt aaa ttc aag gat gag ctt tgg gtt got gtt gca ctg 420
264 lys phe val ser lys phe lys asp glu leu ser val ala val ala leu
265 25               30               35               40
266 aca cta cct gat ggt cat gtt tgg cgt gta gga cta agg aaa got gac 480
268 thr val pro asp gly his val trp arg val gly leu arg lys ala asp
269 45               50               55
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273 asn lys ile trp phe gln asp gly trp gln glu phe val asp arg tyr
274 60               65               70
275 tcc att cgc att ggt cat ctt tgg att ttt aga tat gaa gga aac tct 600
277 ser ile arg ile gly tyr leu leu ile phe arg tyr glu gly asn ser
278 75               80               85
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281 ala phe ser val tyr ile phe asn leu ser his ser glu ile asn tyr
282 90               95               100
283 cat tcc aac ggt ctg atg gat tcc ggt cac aac cac ttc aaa cgc gcc 720
285 his ser thr gly leu met asp ser ala his asn his phe lys arg ala
286 105               110               115               120
287 cgt ttg ttt gaa gac ctt gaa gat gaa gat gcc gag gtc atc ttt cct 780
289 arg leu phe glu asp leu glu asp glu asp ala glu val ile phe pro
290 125               130               135
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293 ser ser val tyr pro ser pro leu pro glu ser thr val pro ala asn
294 140               145               150
295 aaa ggg tat got agt tca gcc atc caa acc ttg ttc act gga cca gtt 900
297 lys gly tyr ala ser ser ala ile gln thr leu phe thr gly pro val
298 155               160               165
299 aca gat gaa gag cca aag cca acc cca aaa ata cct aaa aag aga ggg 960
301 lys ala glu glu pro thr pro thr pro lys ile pro lys lys arg gly
302 170               175               180
303 agg aag aag aaa aat got gat cct gag gaa ata aac tca tca got cgg 1020
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306 185               190               195               200
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**VERIFICATION SUMMARY**

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